SEQUENCE LISTING

```
<110> Friddle, Carl Johan
Hilbun, Erin
Gerhardt, Brenda
Turner, C. Alexander Jr.
```

- <120> Novel Human Ion Channel Protein and Polynucleotides Encoding the Same
- <130> LEX-0251-USA
- <150> US 60/239,623
- <151> 2000-10-10
- <160> 3
- <170> FastSEQ for Windows Version 4.0
- <210> 1
- <211> 1371
- <212> DNA
- <213> homo sapiens
- <400> 1

atggagccgc	ggtgcccgcc	gccgtgcggc	tgctgcgagc	ggctggtgct	caacgtggcc	60	
gggctgcgct	tcgagacgcg	ggcgcgcacg	ctgggccgct	tcccggacac	tctgctaggg	120	
gacccagcgc	gccgcggccg	cttctacgac	gacgcgcgcc	gcgagtattt	cttcgaccgg	180	
caccggccca	gcttcgacgc	cgtgctctac	tactaccagt	ccggtgggcg	gctgcggcgg	240	
ccggcgcacg	tgccgctcga	cgtcttcctg	gaagaggtgg	ccttctacgg	gctgggcgcg	300	
gcggccctgg	cacgcctgcg	cgaggacgag	ggctgcccgg	tgccgcccga	gcgccccctg	360	
ccccgccgcg	ccttcgcccg	ccagctgtgg	ctgcttttcg	agtttcccga	gagctctcag	420	
gccgcgcgcg	tgctcgccgt	agtctccgtg	ctggtcatcc	tcgtctccat	cgtcgtcttc	480	
tgcctcgaga	cgctgcctga	cttccgcgac	gaccgcgacg	gcaċggggct	tgctgctgca	540	
gccgcagccg	gcccgttccc	cgctcggctg	aatggctcca	gccaaatgcc	tggaaatcca	600	
ccccgcctgc	ccttcaatga	cccgttcttc	gtggtggaga	cgctgtgtat	ttgttggttc	660	
tcctttgagc	tgctggtacg	cctcctggtc	tgtccaagca	aggctatctt	cttcaagaac	720	
gtgatgaacc	tcatcgattt	tgtggctatc	cttccctact	ttgtggcact	gggcaccgag	780	
ctggcccggc	agcgaggggt	gggccagcag	gccatgtcac	tggccatcct	gagagtcatc	840	
cgattggtgc	gtgtcttccg	catcttcaag	ctgtcccggc	actcaaaggg	cctgcaaatc	900	
ttgggccaga	cgcttcgggc	ctccatgcgt	gagctgggcc	tcctcatctt	tttcctcttc	960	
atcggtgtgg	tcctctttc	cagcgccgtc	tactttgccg	aagttgaccg	ggtggactcc	1020	
catttcacta	gcatccctga	gtccttctgg	tgggcggtag	tcaccatgac	tacagttggc	1080	
tatggagaca	tggcacccgt	cactgtgggt	ggcaagatag	tgggctctct	gtgtgccatt	1140	
gcgggcgtgc	tgactatttc	cctgccagtg	cccgtcattg	tctccaattt	cagctacttt	1200	
			gggatgttca			1260	
					acctgagcta	1320	
ccacctccac	tctgggcacc	cccagggaaa	cacctggtca	ccgaagtgtg	a	1371	

- <210> 2
- <211> 456
- <212> PRT
- <213> homo sapiens
- <400> 2

Met Glu Pro Arg Cys Pro Pro Pro Cys Gly Cys Cys Glu Arg Leu Val



1				5					10		•			15	
Leu	Asn	Val	Ala 20	Gly	Leu	Arg	Phe	Glu 25	Thr	Arg	Ala	Arg	Thr	Leu	Gly
Arg	Phe	Pro 35	Asp	Thr	Leu	Leu	Gly 40	Asp	Pro	Ala	Arg	Arg 45	Gly	Arg	Phe
Tyr	Asp 50	Asp	Ala	Arg	Arg	Glu 55	Tyr	Phe	Phe	Asp	Arg 60	His	Arg	Pro	Ser
Phe 65	Asp	Ala	Val	Leu	Туr 70	Tyr	Tyr	Gln	Ser	Gly 75	Gly	Arg	Leu	Arg	Arg 80
				85				Phe	90					95	
			100					Arg 105					110		
		115					120	Pro				125			
	130					135		Glu			140				
145					150			Ile		155					160
_				165		_		Arg	170	_			_	175	
			180				_	Pro 185				_	190		_
		195					200	Pro				205			
	210					215		Ile			220				
225					230			Ser		235					240
				245				Ala	250					255	
			260					Arg 265					270		
		275			_		280	Arg			_	285			
	290					295		Gly			300				
305					310			Gly		315					320
				325				Ala	330					335	
			340					11e 345					350		
		355					360	Tyr				365			
	370					375		Leu			380				
385					390			Ile		395					400
				405				Glu	410					415	
			420					Glu 425					430		
		435					440	Pro	Pro	Pro	Leu	Trp 445	Ala	Pro	Pro
GTA	ьуs	Hls	Leu	val	Thr	Glu	val								





1740

1792

450 455

<210> 3 <211> 1792 <212> DNA <213> homo sapiens

<400> 3 60 cggcggcggc cgaggcggcc gaggcggggc cgcaccgggg ccgggcgtcg gggccacacg teggttegeg ggtegeeggg getgegegeg ceatggagee geggtgeeeg eegeegtgeg 120 180 gctgctgcga gcggctggtg ctcaacgtgg ccgggctgcg cttcgagacg cgggcgcgca 240 cgctgggccg cttcccggac actctgctag gggacccagc gcgccgcggc cgcttctacg 300 acgacgcgcg ccgcgagtat ttcttcgacc ggcaccggcc cagcttcgac gccgtgctct 360 actactacca gtccggtggg cggctgcggc ggccggcgca cgtgccgctc gacgtcttcc 420 tggaagaggt ggccttctac gggctgggcg cggcggccct ggcacgcctg cgcgaggacg 480 agggctgccc ggtgccgccc gagcgccccc tgccccgccg cgccttcgcc cgccagctgt ggctgctttt cgagtttccc gagagctctc aggccgcgcg cgtgctcgcc gtagtctccg 540 600 tgctggtcat cctcgtctcc atcgtcgtct tctgcctcga gacgctgcct gacttccgcg 660 acgaccgcga cggcacgggg cttgctgctg cagccgcagc cggcccgttc cccgctcggc 720 tgaatggctc cagccaaatg cctggaaatc cacccgcct gcccttcaat gacccgttct tegtggtgga gaegetgtgt atttgttggt teteetttga getgetggta egeeteetgg 780 840 tctgtccaag caaggctatc ttcttcaaga acgtgatgaa cctcatcgat tttgtggcta 900 tccttcccta ctttgtggca ctgggcaccg agctggcccg gcagcgaggg gtgggccagc 960 aggecatgte actggecate etgagagtea teegattggt gegtgtette egeatettea 1020 agetyteecy geacteaaay ggeetyeaaa tettyggeea gaegettegy geeteeatye 1080 gtgagctggg cctcctcatc tttttcctct tcatcggtgt ggtcctcttt tccagcgccg 1140 tctactttgc cgaagttgac cgggtggact cccatttcac tagcatccct gagtccttct ggtgggcggt agtcaccatg actacagttg gctatggaga catggcaccc gtcactgtgg 1200 1260 gtggcaagat agtgggctct ctgtgtgcca ttgcgggcgt gctgactatt tccctgccag 1320 tgcccgtcat tgtctccaat ttcagctact tttatcaccg ggagacagag ggcgaagagg ctgggatgtt cagccatgtg gacatgcagc cttgtggccc actggagggc aaggccaatg 1380 1440 gggggctggt ggacggggag gtacctgagc taccacctcc actctgggca cccccaggga 1500 aacacctggt caccgaagtg tgaggaacag ttgaggtctg caggacctca cacctcccta 1560 gagggaggga gggagggcag ggtggagggc aaggctgggg ggaggggatt gggtttagga 1620 agagctaggt taagtcrtaa cgagtgggga aacactgagt cttgttgggt cttgggttgt 1680 gtggtttggt agctcctgtg ggtacctcct gaagcagcag cgaatggcaa tgggttgtgt

tgtgttaatg aagactcaat tggttcatat tactctgagt tgtgcaaagc tcatggagcc

ttttggggta gtgttgagat aggtttggtc rtatcatttt gtgagtttcc ta